



## SEQUENCE LISTING

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<120> CONJUGATES COMPRISING HUMAN IL-18 AND
SUBSTITUTION MUTANTS THEREOF

<130> PU60053

<140> 10/823,964

<141> 2004-04-14

<150> 60/462,947

<151> 2003-04-15

<160> 28

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 157

<212> PRT

<213> Homo sapiens

<400> 1

Tyr Phe Gly Lys Leu Glu Ser Lys Leu Ser Val Ile Arg Asn Leu Asn 1 5 10 15

1 5 10 15
Asp Gln Val Leu Phe Ile Asp Gln Gly Asn Arg Pro Leu Phe Glu Asp

20 25 30 Met Thr Asp Ser Asp Cys Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile

35 40 45

Ile Ser Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile

50 55 60
Ser Val Lys Cys Glu Lys Ile Ser Thr Leu Ser Cys Glu Asn Lys Ile

65 70 75 80
Ile Ser Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp Thr Lys

85 90 95 Ser Asp Ile Ile Phe Phe Gln Arg Ser Val Pro Gly His Asp Asn Lys

Met Gln Phe Glu Ser Ser Ser Tyr Glu Gly Tyr Phe Leu Ala Cys Glu
115 120 125

Lys Glu Arg Asp Leu Phe Lys Leu Ile Leu Lys Lys Glu Asp Glu Leu 130 135 140

Gly Asp Arg Ser Ile Met Phe Thr Val Gln Asn Glu Asp 145 150

<210> 2

<211> 157

<212> PRT <213> Mus musculus

<400> 2 Asn Phe Gly Arg Leu His Cys Thr Thr Ala Val Ile Arg Asn Ile Asn Asp Gln Val Leu Phe Val Asp Lys Arg Gln Pro Val Phe Glu Asp Met 25 Thr Asp Ile Asp Gln Ser Ala Ser Glu Pro Gln Thr Arg Leu Ile Ile Tyr Met Tyr Lys Asp Ser Glu Val Arg Gly Leu Ala Val Thr Leu Ser 55 Val Lys Asp Ser Lys Met Ser Thr Leu Ser Cys Lys Asn Lys Ile Ile 75 70 Ser Phe Glu Glu Met Asp Pro Pro Glu Asn Ile Asp Asp Ile Gln Ser 90 Asp Leu Ile Phe Phe Gln Lys Arg Val Pro Gly His Asn Lys Met Glu 105 Phe Glu Ser Ser Leu Tyr Glu Gly His Phe Leu Ala Cys Gln Lys Glu 120 Asp Asp Ala Phe Lys Leu Ile Leu Lys Lys Lys Asp Glu Asn Gly Asp 135 Lys Ser Val Met Phe Thr Leu Thr Asn Leu His Gln Ser

<210> 3 <211> 203 <212> PRT <213> Homo sapiens

<400> 3

Met His His His His His Thr Arg Gly Met Ala Ala Glu Pro Val 10 Glu Asp Asn Cys Ile Asn Phe Val Ala Met Lys Phe Ile Asp Asn Thr 20 Leu Tyr Phe Ile Ala Glu Asp Asp Glu Asn Leu Glu Ser Asp Tyr Phe 40 Gly Lys Leu Glu Ser Lys Leu Ser Val Ile Arg Asn Leu Asn Asp Gln 55 Val Leu Phe Ile Asp Gln Gly Asn Arg Pro Leu Phe Glu Asp Met Thr 75 Asp Ser Asp Cys Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile Ile Ser 90 Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile Ser Val 105 Lys Cys Glu Lys Ile Ser Thr Leu Ser Cys Glu Asn Lys Ile Ile Ser 120 125 Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp Thr Lys Ser Asp Ile Ile Phe Phe Gln Arg Ser Val Pro Gly His Asp Asn Lys Met Gln 150 155 Phe Glu Ser Ser Tyr Glu Gly Tyr Phe Leu Ala Cys Glu Lys Glu 165 170 Arg Asp Leu Phe Lys Leu Ile Leu Lys Lys Glu Asp Glu Leu Gly Asp

185

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Arg Ser Ile Met Phe Thr Val Gln Asn Glu Asp
   195
<210> 4
<211> 157
<212> PRT
<213> Homo sapiens
<220>
<223> Whereby the Cysteine at position 38 of this human IL-18
      sequence has been replaced with Serine.
<400> 4
Tyr Phe Gly Lys Leu Glu Ser Lys Leu Ser Val Ile Arg Asn Leu Asn
                                    10
Asp Gln Val Leu Phe Ile Asp Gln Gly Asn Arg Pro Leu Phe Glu Asp
            20
                                25
Met Thr Asp Ser Asp Ser Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile
        35
                            40
Ile Ser Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile
                        55
Ser Val Lys Cys Glu Lys Ile Ser Thr Leu Ser Cys Glu Asn Lys Ile
                    70
                                        75
Ile Ser Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp Thr Lys
                                    90
Ser Asp Ile Ile Phe Phe Gln Arg Ser Val Pro Gly His Asp Asn Lys
                                105
                                                     110
Met Gln Phe Glu Ser Ser Ser Tyr Glu Gly Tyr Phe Leu Ala Cys Glu
                            120
                                                125
Lys Glu Arg Asp Leu Phe Lys Leu Ile Leu Lys Lys Glu Asp Glu Leu
                        135
Gly Asp Arg Ser Ile Met Phe Thr Val Gln Asn Glu Asp
145
                    150
<210> 5
<211> 157
<212> PRT
<213> Homo sapiens
<220>
<223> Whereby the Cysteine at position 38 of this human IL-18
      sequence has been replaced with Serine, the Cysteine at
      position 68 has been replaced with Aspartic acid, and the
      Asparagine at position 78 has been replaced with Cysteine.
<400> 5
Tyr Phe Gly Lys Leu Glu Ser Lys Leu Ser Val Ile Arg Asn Leu Asn
                                    1.0
Asp Gln Val Leu Phe Ile Asp Gln Gly Asn Arg Pro Leu Phe Glu Asp
            20
                                25
Met Thr Asp Ser Asp Ser Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile
```

40 Ile Ser Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile

35

50

```
Ser Val Lys Asp Glu Lys Ile Ser Thr Leu Ser Cys Glu Cys Lys Ile
                    70
Ile Ser Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp Thr Lys
                85
                                    90.
Ser Asp Ile Ile Phe Phe Gln Arg Ser Val Pro Gly His Asp Asn Lys
            100
                                105
Met Gln Phe Glu Ser Ser Tyr Glu Gly Tyr Phe Leu Ala Cys Glu
        115
                            120
                                                125
Lys Glu Arg Asp Leu Phe Lys Leu Ile Leu Lys Lys Glu Asp Glu Leu
                        135
Gly Asp Arg Ser Ile Met Phe Thr Val Gln Asn Glu Asp
                    150
```

<210> 6 <211> 157 <212> PRT <213> Homo sapiens

<220>

<223> Whereby the Cysteine at position 38 of thi human IL-18 sequence has been replaced with Serine, the Cysteine at position 68 has been replaced with Aspartic acid, and the Glutamic acid at position 121 has been replaced with Cysteine.

<400> 6 Tyr Phe Gly Lys Leu Glu Ser Lys Leu Ser Val Ile Arg Asn Leu Asn 10 Asp Gln Val Leu Phe Ile Asp Gln Gly Asn Arg Pro Leu Phe Glu Asp 25 Met Thr Asp Ser Asp Ser Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile 45 Ile Ser Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile Ser Val Lys Asp Glu Lys Ile Ser Thr Leu Ser Cys Glu Asn Lys Ile 70 Ile Ser Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp Thr Lys 90 Ser Asp Ile Ile Phe Phe Gln Arg Ser Val Pro Gly His Asp Asn Lys 100 105 Met Gln Phe Glu Ser Ser Ser Tyr Cys Gly Tyr Phe Leu Ala Cys Glu 120 125 Lys Glu Arg Asp Leu Phe Lys Leu Ile Leu Lys Lys Glu Asp Glu Leu 135 Gly Asp Arg Ser Ile Met Phe Thr Val Gln Asn Glu Asp

<210> 7 <211> 157 <212> PRT <213> Homo sapeins

<220>

<223> Whereby the Cysteine at position 38 of this human IL-18 sequence has been replaced with Serine, the Cysteine at position 68 has been replaced with Aspartic acid, and the Leucine at position 144

has been replaced with Cysteine.

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<400> 7
Tyr Phe Gly Lys Leu Glu Ser Lys Leu Ser Val Ile Arg Asn Leu Asn
Asp Gln Val Leu Phe Ile Asp Gln Gly Asn Arg Pro Leu Phe Glu Asp
Met Thr Asp Ser Asp Ser Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile
                            40
Ile Ser Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile
                        55--
Ser Val Lys Asp Glu Lys Ile Ser Thr Leu Ser Cys Glu Asn Lys Ile
                    70
                                        75
Ile Ser Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp Thr Lys
                85
                                    90
Ser Asp Ile Ile Phe Phe Gln Arg Ser Val Pro Gly His Asp Asn Lys
                                105
Met Gln Phe Glu Ser Ser Tyr Glu Gly Tyr Phe Leu Ala Cys Glu
                            120
                                                125
Lys Glu Arg Asp Leu Phe Lys Leu Ile Leu Lys Lys Glu Asp Glu Cys
                        135
Gly Asp Arg Ser Ile Met Phe Thr Val Gln Asn Glu Asp
145
                    150
```

```
<210> 8
<211> 157
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<220>

<223> Whereby the Cysteine at position 38 of the human IL-18 sequence has been replaced with Serine, the Cysteine at position 68 has been replaced with Aspartic acid, and Aspartic acid at position 157 has been replaced with Cysteine.

```
<400> 8
Tyr Phe Gly Lys Leu Glu Ser Lys Leu Ser Val Ile Arg Asn Leu Asn
Asp Gln Val Leu Phe Ile Asp Gln Gly Asn Arg Pro Leu Phe Glu Asp
                                25
Met Thr Asp Ser Asp Ser Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile
Ile Ser Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile
                        55
Ser Val Lys Asp Glu Lys Ile Ser Thr Leu Ser Cys Glu Asn Lys Ile
                    70
                                        75
Ile Ser Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp Thr Lys
                                    90
Ser Asp Ile Ile Phe Phe Gln Arg Ser Val Pro Gly His Asp Asn Lys
            100
                                105
                                                    110
Met Gln Phe Glu Ser Ser Tyr Glu Gly Tyr Phe Leu Ala Cys Glu
        115
                            120
                                                125
Lys Glu Arg Asp Leu Phe Lys Leu Ile Leu Lys Lys Glu Asp Glu Leu
   130
                        135
Gly Asp Arg Ser Ile Met Phe Thr Val Gln Asn Glu Cys
                    150
```

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo sapiens

```
<210> 9
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<211> 157

<212> PRT

<213> Homo sapeins

<220>

<223> Whereby the Cysteine at position 38 of the human IL-18 sequence has been replaced with Serine, the Cysteine at position 68 has been replaced with Serine, and Leucine at position 144 has been replaced with Cysteine.

<400> 9
Tyr Phe Gly Lys Leu Glu Ser Lys Leu Ser Val Ile Arg Asn Leu Asn

1 5 10 15
Asp Gln Val Leu'Phe Ile Asp Gln Gly Asn Arg Pro Leu Phe Glu Asp

Met Thr Asp Ser Asp Ser Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile 35 40 45

Ile Ser Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile

Ser Val Lys Ser Glu Lys Ile Ser Thr Leu Ser Cys Glu Asn Lys Ile
65 70 75 80

Ile Ser Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp Thr Lys
85 90 95

Ser Asp Ile Ile Phe Phe Gln Arg Ser Val Pro Gly His Asp Asn Lys 100 105 110

Met Gln Phe Glu Ser Ser Tyr Glu Gly Tyr Phe Leu Ala Cys Glu 115 120 125

Lys Glu Arg Asp Leu Phe Lys Leu Ile Leu Lys Lys Glu Asp Glu Cys
130 135 140

Gly Asp Arg Ser Ile Met Phe Thr Val Gln Asn Glu Asp 145 150 155

<210> 10

<211> 157

<212> PRT

<213> Homo sapiens

<220>

<223> Whereby the Cysteine at position 38 of the human IL-18 sequence has been replaced with Serine, the Cysteine at position 68 has been replaced with Serine, and Aspartic acid at position 157 has been replaced with Cysteine.

<400> 10

Tyr Phe Gly Lys Leu Glu Ser Lys Leu Ser Val Ile Arg Asn Leu Asn 1 5 10 15

Asp Gln Val Leu Phe Ile Asp Gln Gly Asn Arg Pro Leu Phe Glu Asp 20 25 30

Met Thr Asp Ser Asp Ser Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile
35 40 45

```
Ile Ser Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile
    50 ·
                        55
Ser Val Lys Ser Glu Lys Ile Ser Thr Leu Ser Cys Glu Asn Lys Ile
                    70
                                        75
Ile Ser Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp Thr Lys
                                    90
Ser Asp Ile Ile Phe Phe Gln Arg Ser Val Pro Gly His Asp Asn Lys
            100
                                105
                                                     110
Met Gln Phe Glu Ser Ser Tyr Glu Gly Tyr Phe Leu Ala Cys Glu
       115
                            120
                                                125
Lys Glu Arg Asp Leu Phe Lys Leu Ile Leu Lys Lys Glu Asp Glu Leu
                       135 .
Gly Asp Arg Ser Ile Met Phe Thr Val Gln Asn Glu Cys
                    150
<210> 11
<211> 4
<212> PRT
<213> Artificial Sequence
<220>
<223> Tryptic peptides predicted for S-carboxymethylated
      wild type IL-18
<400> 11
Tyr Phe Gly Lys
<210> 12
<211> 4
<212> PRT
<213> Artificial Sequence
<220>
<223> Tryptic peptides predicted for S-carboxymethylated
      wild type IL-18
<400> 12
Leu Glu Ser Lys
<210> 13
<211> 5
<212> PRT
<213> Artificial Sequence
<220>
<223> Tryptic peptides predicted for S-carboxymethylated
      wild type IL-18
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<400> 13

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Leu Ser Val Ile Arg
1 . 5
<210> 14
<211> 26
<212> PRT
<213> Artificial Sequence
<220>
<223> Tryptic peptides predicted for S-carboxymethylated
      wild type IL-18
Asn Leu Asn Asp Gln Val Leu Phe Ile Asp Gln Gly Asn Arg Pro Leu
               5
                                   10
Phe Glu Asp Met Thr Asp Ser Asp Cys Arg
<210> 15
<211> 5
<212> PRT
<213> Artificial Sequence
<220>
<223> Tryptic peptides predicted for S-carboxymethylated
      wild type IL-18
<400> 15
Asp Asn Ala Pro Arg
1
                5
<210> 16
<211> 9
<212> PRT
<213> Artificial Sequence ·
<220>
<223> Tryptic peptides predicted for S-carboxymethylated
     wild type IL-18
<400> 16
Thr Ile Phe Ile Ile Ser Met Tyr Lys
<210> 17
<211> 5
<212> PRT
<213> Artificial Sequence
<220>
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<223> Tryptic peptides predicted for S-carboxymethylated
      wild type IL-18
<400> 17
Asp Ser Gln Pro Arg
<210> 18
<211> 9
<212> PRT
<213> Artificial Sequence
<223> Tryptic peptides predicted for S-carboxymethylated
      wild type IL-18
<400> 18
Gly Met Ala Val Thr Ile Ser Val Lys
   5
<210> 19
<211> 9
<212> PRT
<213> Artificial Sequence
<220>
<223> Tryptic peptides predicted for S-carboxymethylated
      wild type IL-18
<400> 19
Ile Ser Thr Leu Ser Cys Glu Asn Lys
                5
<210> 20
<211> 5
<212> PRT
<213> Artificial Sequence
<220>
<223> Tryptic peptides predicted for S-carboxymethylated
     wild type IL-18
<400> 20
Ile Ile Ser Phe Lys
<210> 21
<211> 9
<212> PRT
```

```
<213> Artificial Sequence
<220>
<223> Tryptic peptides predicted for S-carboxymethylated
      wild type IL-18
<400> 21
Glu Met Asn Pro Pro Asp Asn Ile Lys
<210> 22
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Tryptic peptides predicted for S-carboxymethylated
      wild type IL-18
<400> 22
Ser Asp Ile Ile Phe Phe Gln Arg
<210> 23
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Tryptic peptides predicted for S-carboxymethylated
      wild type IL-18
<400> 23
Ser Val Pro Gly His Asp Asn Lys
 1
                 5
<210> 24
<211> 17
<212> PRT
<213> Artificial Sequence
<220>
<223> Tryptic peptides predicted for S-carboxymethylated
      wild type IL-18
<400> 24
Met Gln Phe Glu Ser Ser Ser Tyr Glu Gly Tyr Phe Leu Ala Cys Glu
1
                                    10
                                                         15
Lys
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<210> 25
<211> 4
<212> PRT
<213> Artificial Sequence
<220>
<223> Tryptic peptides predicted for S-carboxymethylated
      wild type IL-18
<400> 25
Asp Leu Phe Lys
<210> 26
<211> 4
<212> PRT
<213> Artificial Sequence
<220>
<223> Tryptic peptides predicted for S-carboxymethylated
      wild type IL-18
<400> 26
Leu Ile Leu Lys
 1
<210> 27
<211> 7
<212> PRT
<213> Artificial Sequence
<220>
<223> Tryptic peptides predicted for S-carboxymethylated
      wild type IL-18
<400> 27
Glu Asp Glu Leu Gly Asp Arg
<210> 28
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Tryptic peptides predicted for S-carboxymethylated
     wild type IL-18
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